RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/550,410
Source:	IFWP.
Date Processed by STIC:	. 1/9/07

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IFWP

RAW SEQUENCE LISTING DATE: 01/09/2007 PATENT APPLICATION: US/10/550,410 TIME: 09:35:43

Input Set : A:\3022-109.ST25.txt

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3 <110> APPLICANT: Burnie, James
         Matthews, Ruth C.
 4
 6 <120> TITLE OF INVENTION: Treatment of Infection Due to Clostridium Difficile
 8 <130> FILE REFERENCE: 3022-109
10 <140> CURRENT APPLICATION NUMBER: 10/550,410
11 <141> CURRENT FILING DATE: 2005-09-23
13 <150> PRIOR APPLICATION NUMBER: PCT/GB2004/001383
14 <151> PRIOR FILING DATE: 2004-03-25
16 <150> PRIOR APPLICATION NUMBER: GB0306782.4
17 <151> PRIOR FILING DATE: 2003-03-25
19 <160> NUMBER OF SEQ ID NOS: 8
21 <170> SOFTWARE: PatentIn version 3.3
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 975
25 <212> TYPE: DNA
26 <213> ORGANISM: Clostridium difficile
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30 <221> NAME/KEY: CDS
31 <222> LOCATION: (40)..(972)
33 <400> SEQUENCE: 1
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                                                                          54
35
                                              Met Lys Ile Leu Val
36
38 ttt gga gca cgc gat tat gaa gaa cca gta ata aaa aaa tgg tct gaa
                                                                         102
39 Phe Gly Ala Arg Asp Tyr Glu Glu Pro Val Ile Lys Lys Trp Ser Glu
42 gaa cat aag gat gtt caa gtg gat att tat cct gaa aac atg act gaa
                                                                         150
43 Glu His Lys Asp Val Gln Val Asp Ile Tyr Pro Glu Asn Met Thr Glu
46 gaa aat gta gtt aaa gct aaa ggg tat gat ggt ata tct ata caa caa
                                                                         198
47 Glu Asn Val Val Lys Ala Lys Gly Tyr Asp Gly Ile Ser Ile Gln Gln
48
           40
                               45
50 act aac tat ata gat aat cct tat att tat gaa act tta aaa gat gct
                                                                         246
51 Thr Asn Tyr Ile Asp Asn Pro Tyr Ile Tyr Glu Thr Leu Lys Asp Ala
                                               65
54 ggg gtt aaa gtt ata gct tca aga act gca ggg gtt gac atg ata cat
                                                                         294
55 Gly Val Lys Val Ile Ala Ser Arg Thr Ala Gly Val Asp Met Ile His
                       75
                                           80
58 ttt gat tta gtt aat gaa aat gga ctt atc gtt aca aac gtt cct gct
                                                                         342
59 Phe Asp Leu Val Asn Glu Asn Gly Leu Ile Val Thr Asn Val Pro Ala
                                       95
62 tat tca cct aat gca ata gct gaa tta gct gtt act caa gct atg aac
                                                                         390
63 Tyr Ser Pro Asn Ala Ile Ala Glu Leu Ala Val Thr Gln Ala Met Asn
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			aaa aaa gtc tgt Lys Lys Val Cys	
		125	130	Gid Giy Asp
	=			tat att acc 100
			aca gaa gtt aga	
•	p lie Ala Giu	-	Thr Glu Val Arg	ser lie inr
72 135		140	145	534
			ggt gct act tct	
_	-		Gly Ala Thr Ser	_
76 150	155		160	165
		_	gca tat gac caa	
	=	Asn val lie	Ala Tyr Asp Gln	_
80	170		175	180
			aaa gat tct tta	
-	-	-	Lys Asp Ser Leu	-
84	185	190		195
_			cat act cct tta	
_	_		His Thr Pro Leu	Leu Glu Gly
88 20		205	210	
		-	cta gct ata atg	
_	s Met Ile Asn		Leu Ala Ile Met	Lys Asp Gly
92 215		220	225	
			ggt tta att gat	
			Gly Leu Ile Asp	
96 230	235		240	245
=			att aga gct gct	-
	_	Ser Gly Lys	Ile Arg Ala Ala	_
100	250		255	260
-		-	aac aaa aaa atg	
		_	Asn Lys Lys Met	-
104	265	270		275
			ctt ctt tct ato	
			: Leu Leu Ser Met	
	30	285	290	
			act agt aca gco	
	nr His His Le	=	Thr Ser Thr Ala	a lle Glu Asn
112 295		300	305	
114 ata gtt ta	aa			975
115 Ile Val				
116 310				
119 <210> SEQ				
120 <211> LENG				
121 <212> TYPI			-	•
122 <213> ORGZ		iaium diffici	.ıe	
124 <400> SEQU		a3		
		e Giy Ala Arg	Asp Tyr Glu Glu	
127 1	5		10	15
			Val Gln Val Asp	
131	20	25		30

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134 135	Glu	Asn	Met 35	Thr	Glu	Glu	Asn	Val	Val	Lys	Ala	Lys	Gly 45	Tyr	Asp	Gly	
138	Ile			Gln	Gln	Thr			Ile	Asp	Asn			Ile	Tyr	Glu	
139	Mla sa	50	T	7.00	7.7.	a 1	55	T	77-7	T1.	77.	60	7 ~~~	The	ח ד ה	C1	
142		Leu	ьys	Asp	Ala	70	vai	гуѕ	vai	TIE	75	ser	Arg	Inr	Ald	80	
		Asp	Met	Ile	His		Asp	Leu	Val	Asn		Asn	Gly	Leu	Ile		
147		-			85		_			90			_		95		
	Thr	Asn	Val		Ala	Tyr	Ser	Pro		Ala	Ile	Ala	Glu		Ala	Val	
151	1	~3		100		_			105		_	_	**- 7	110		•	
154	Thr	Gin	115	Met	Asn	Leu	Leu	Arg	ьуs	Thr	Pro	Leu	vai 125	ьуs	ьуs	ьуs	
	Val	Cvs		Glv	Asp	Tvr	Ara		Tle	Ala	Glu	Leu		Glv	Thr	Glu	
159	· · · ·	130	OLU	ردن	n.sp	-1-	135	115			014	140	LCu	017		014	
162	Val	Arg	Ser	Ile	Thr	Val	Gly	Val	Ile	Gly	Thr	Gly	Lys	Ile	Gly	Ala	
	145					150			•		155					160	
	Thr	Ser	Ala	Lys	Leu	Phe	Lys	Gly	Leu	_	Ala	Asn	Val	Ile		Tyr	
167	_	~7	_	_	165	_	_	_	_	170		_	m1 .	.	175	•	
170	Asp	Gin	Tyr	180	Asn	ser	Asp	Leu	185	Asp	тте	Leu	Thr	191	гÀг	Asp	
	Sor	T.OU	Glu		Leu	T.211	Larg	Glu		λαη	Val	Tla	Thr		Hie	Thr	
175	Ser	Deu	195	лор	пец	пец	цуз	200	AIG	App	Vai	116	205	пси	1115	*111	
	Pro	Leu		Glu	Gly	Thr	Lys		Met	Ile	Asn	Lys		Thr	Leu	Ala	
179		210			-		215					220	-				
182	Ile	Met	Lys	Asp	Gly	Ala	Tyr	Ile	Val	Asn	Thr	Ala	Arg	Gly	Gly	Leu	
	225					230					235					240	
	Ile	Asp	Thr	Gly	Asp	Leu	Ile	Glu	Ala		Asp	Ser	Gly	Lys		Arg	
187	77.	77.	77.	T	245	ml	nh -	a1	mla aa	250	0 1	T	Dha	T	255	T	
190	Ата	Ата	Ата	260	Asp	Thr	Pne	GIU	265	GIU	GIY	Leu	Pne	270	ASII	ьуѕ	
	Lvs	Met	Asn		Gly	Glu	Leu	Thr		Pro	Glu	Tle	Asn		Leu	Leu	
195	-,-		275		J-1			280	P				285	-1-			
	Ser	Met		Gln	Val	Ile	Phe		His	His	Leu	Gly	Phe	Phe	Thr	Ser	
199		290					295					300					
202	Thr	Ala	Ile	Glu	Asn	Ile	Val										
203						310											
			EQ II														
	_	_	ENGTI		0												
			YPE:		01 - 1		a 2	2: 6		1 -							
			EQUEI		Clos	SCIIC	11 UIII	alli	LICI.	re							
			-			aa											20
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Input Set : A:\3022-109.ST25.txt

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254 <213> ORGANISM: Clostridi	ium difficile
256 <400> SEQUENCE: 8	
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259 1 5	10

VERIFICATION SUMMARY

DATE: 01/09/2007

PATENT APPLICATION: US/10/550,410

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